## SEQUENCE LISTING

<110> Allen, Keith D.

<120> TRANSGENIC MICE CONTAINING CALCIUM ION
 CHANNEL (Trp6) GENE DISRUPTIONS

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<151> 2001-03-29
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Ile Asn Ala Tyr Lys Gly Leu Ala Ser Pro Ala Tyr Leu Ser Leu Ser Ser Glu Asp Pro Val Met Thr Ala Leu Glu Leu Ser Asn Glu Leu Ala Val Leu Ala Asn Ile Glu Lys Glu Phe Lys Asn Asp Tyr Arg Lys Leu Ser Met Gln Cys Lys Asp Phe Val Val Gly Leu Leu Asp Leu Cys Arg Asn Thr Glu Glu Val Glu Ala Ile Leu Asn Gly Asp Ala Glu Thr Arg Gin Pro Gly Asp Phe Gly Arg Pro Asn Leu Ser Arg Leu Lys Leu Ala Ile Lys Asp Glu Val Lys Lys Phe Val Ala His Pro Asn Cys Gln Gln Gln Leu Leu Ser Ile Trp Tyr Glu Asn Leu Ser Gly Leu Arg Gln Gln Thr Met Ala Val Lys Phe Leu Val Val Leu Ala Val Ala Ile Gly Leu Pro Phe Leu Ala Leu Ile Tyr Trp Cys Ala Pro Cys Ser Lys Met Gly Lys Ile Leu Pro Arg Pro Phe Met Lys Phe Val Ala His Ala Ala Ser Phe Thr Ile Phe Leu Gly Leu Leu Val Met Asn Ala Ala Asp Arg Phe Glu Gly Thr Lys Leu Leu Pro Asn Glu Thr Ser Thr Asp Asn Ala Arg Gln Leu Phe Arg Met Lys Thr Ser Cys Phe Ser Trp Met Glu Met Leu Ile Ile Ser Trp Val Ile Gly Met Ile Trp Ala Glu Cys Lys Glu Ile Trp Thr Gln Gly Pro Lys Glu Tyr Leu Phe Glu Leu Trp Asn Met Leu Asp Phe Gly Met Leu Ala Ile Phe Ala Ala Ser Phe Ile Ala Arg Phe Met Ala Phe Trp His Ala Ser Lys Ala Gln Ser Ile Ile Asp Ala Asn Asp Thr Leu Lys Asp Leu Thr Lys Val Thr Leu Gly Asp Asn Val Lys Tyr Tyr Asn Leu Ala Arg Ile Lys Trp Asp Pro Thr Asp Pro Gln Ile Ile Ser Glu Gly Leu Tyr Ala Ile Ala Val Val Leu Ser Phe Ser Arg Ile Ala Tyr Ile Leu Pro Ala Asn Glu Ser Phe Gly Pro Leu Gln Ile Ser Leu Gly Arg Thr Val Lys Asp Ile Phe Lys Phe Met Val Ile Phe Ile Met Val Phe Val Ala Phe Met Ile Gly Met Phe Asn Leu Tyr Ser Tyr Tyr Ile Gly Ala Lys Gln Asn Glu Ala Phe Thr Thr Val Glu Glu Ser Phe Lys Thr Leu Phe Trp Ala Ile Phe Gly Leu Ser Glu Val Lys Ser Val Val Ile Asn Tyr Asn His Lys Phe Ile Glu Asn Ile Gly Tyr Val Leu Tyr Gly Val Tyr Asn Val Thr Met Val Ile Val Leu Leu Asn 7 Ú 5 Met Leu Ile Ala Met Ile Asn Ser Ser Phe Gln Glu Ile Glu Asp Asp Ala Asp Val Glu Trp Lys Phe Ala Arg Ala Lys Leu Trp Phe Ser Tyr 

Phe Glu Glu Gly Arg Thr Leu Pro Val Pro Phe Asn Leu Val Pro Ser 755 760 Pro Lys Ser Leu Leu Tyr Leu Leu Leu Lys Phe Lys Lys Trp Met Cys 770 775 780 Glu Leu Ile Gln Gly Gln Lys Gln Gly Phe Gln Glu Asp Ala Glu Met 785 790 795 Asn Lys Arg Asn Glu Glu Lys Lys Phe Gly Ile Ser Gly Ser His Glu 805 810 Asp Leu Ser Lys Phe Ser Leu Asp Lys Asn Gln Leu Ala His Asn Lys 820 825 830 Gln Ser Ser Thr Arg Ser Ser Glu Asp Tyr His Leu Asn Ser Phe Ser 840 835 845 Asn Pro Pro Arg Gln Tyr Gln Lys Ile Met Lys Arg Leu Ile Lys Arg 850 855 860 Tyr Val Leu Gln Ala Gln Ile Asp Lys Glu Ser Asp Glu Val Asn Glu 865 870 875 880 Gly Glu Leu Lys Glu Ile Lys Gln Asp Ile Ser Ser Leu Arg Tyr Glu 890 885 Leu Leu Glu Glu Lys Ser Gln Asn Ser Glu Asp Leu Ala Glu Leu Ile 900 905 Arg Lys Leu Gly Glu Arg Leu Ser Leu Glu Pro Lys Leu Glu Glu Ser 915 920 925 Arg Arg 930 <210> 3 <211> 200 <212> DNA <213> Artificial Sequence <220> <223> Targeting vector <400> 3 tecteaatte taactgeatt tettetggaa aagaataaaa egatteacea gageteeaga 60 ggatageeta agetgagttg tttttaatea aateattetg tgtgetgtet eacceetagt 120 ttgtggctca tccaagctgt cagcaacagc tcctgtccat atggtatgag aacctctctg 180 gtttacggca gcagaccatg

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